



FIG. 1A

FIG. 1A
FIG. 1B
FIG. 1C

FIG. 1

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1 CCCTTCTCAGGACTCTGGCTGCCAGCAGCTCCGCCCTTTCAGATCAATTCTCGACCACC 60
61 CACCTTGGGACTGCCGCCAGTCCCTGCTGCCCTCTGGATCAGTGGGTCCAGACACGCCCCCT 120
121 CCAGGACCTCAAAGCACCCCGACCTAAGGTACACAGCCCACTGGCCCCAGACGCA GTGG 180
181 GCTCCGCTGACTCTCTTGGACACCTCCTGGAGAGAAATGCTCCTGTCTGCCATCGTTT 240
M L P V C H R P
241 TTGCGACCACCTCCTCCTCCTGCTCTTGCTGCCCTCGACGACCCCTGGCCCCCGGCCAGC 300
C A H L L L L L L L P S T T L A P A P A
301 ATCCATGGGCCCGCTGCCGCCCTGCTCCAGGTTCTTGGGCTTCCCGAAGCGCCCCGGAG 360
S M G P A A A L L G V L G L P E A P R S
361 CGTCCCCACACCGACCTGTGCTCCTCCTGTATGTGGCGCCTATTCCGTCGCCGTGACCC 420
V P T H R P P V P P V M W R L P A A A P
421 CCAGGAGGCCAGAGTGGGACGCCCTCTGCGGCCATGCCACGTGGAGGAAC TAGGGTCCG 480
Q E A R V G R P L R P C H V E E L G V A
481 CGGAAACATTGTGCGCCACATCCCCGACAGCGGTCTGTCTCCAGGCCCGCACAAACCCGC 540
G N I V R R H I P D S G L S S R P A Q P A



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541 CAGGACCTCGGGGCTGTGCCCCGAGTGACAGTCGTCTTTGACCTGTGCAATGTGGAGCC 600
R T S G L C P E W T V F D L S N V E P

601 CACAGAGCGCCCAACACGCGCGCTTAGACTTGGCGCTGGAGGCTGAGTGTGAAGATAC 660
T E R P T R A R L E L R L E A E C E D T

661 AGGAGGGTGGGAGCTAAGCGTGGCACTGTGGGCCGACGCAGAGCATCCAGGGCCTGAGCT 720
G G W E L S V A L W A D A E H P G P E L

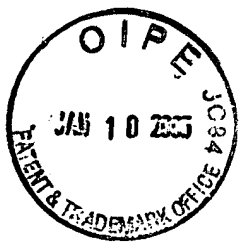
721 GCTGCGCGTGGCGGCCACACAGGGGTGCTCCTGCGCGCAGACCTACTGGGGACTGCAGT 780
L R V P A P P G V L L R A D L L G T A V

781 AGCCGCCAACGCATCAGTGCCCTGTACTGTGCGCCTGGCGCTGTCACTGCACCCCTGGGCG 840
A A N A S V P C T V R L A L S L H P G A

841 CACTGCAGCCTGTGGCGCCTGGCTGAGGCCCTCCCTGCTGCTGTTGACGCTGGACCCACG 900
T A A C G R L A E A S L L L V T L D P R

901 CCTGTGTCCCTTGCCGCGATGCGGCGCCACACGAGCCAGGGTAGAAGTTGGTCCAGT 960
L C P L P R L R R H T E P R V E V G P V

FIG. 1B



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961 GGGCACTTGTCTACCCGACGGTTGCATGTGAGCTTCCGTGAGGTGGGCTGGCACCGTTG 1020
G T C R T R R L H V S F R E V G W H R W

1021 GGTGATCGCGCGGTGGCTTCCTAGCCAACTTCTGCCAGGGCACGTGCGCACTACCCGA 1080
V I A P R G G F L A N F C Q G T C A L P E

1081 AACGCTGAGGGACCCGGCGGCGCTGCACTCAACCACGCTGTGCTGCGCGCTCAT 1140
T L R G P G G P P A L N H A V L R A L M

1141 GCACGCAGCTGTCTCCACCCGGGTGCAGGCTGCCCTGCTGCGTGCCAGAGCGTCTATC 1200
H A A A P T P G A G S P C C V P E R L S

1201 ACCCATCTCCGTGCTCTTCTTCGACAATAGTGACAACGTGGTCTCGCACACTACGAAGA 1260
P I S V L F F D N S D N V V L R H Y E D

1261 CATGGTGGTGATGAGTGTGGCTGCCGTTGACCACCCGGGACACCCCTTTCAGGGACCGCC 1320
M V V D E C G C R

1321 CCACGCAAAAGCAGGGACTGTTTGTTCATGTTTATTGTTGACAAAAGCTTAAACAAA 1380
1381 TTTGACT 1387

FIG. 1C